

INTERPRETATION TABLE								
DRB1*09 SSP subtyping								
Amplification patterns of the DRB1*09 alleles								
	Well ⁵							
	1	2	3	4	5	6	7	8
Length of spec.	195	130	185	215	185	250	220	175
PCR product					215	275		
Length of int.	515	430	430	430	430	430	430	430
pos. control ¹								
5'-primer ²	26(164)	28(170)	9(112)	9(112)	9(112)	9(112)	26(165)	13(125)
	5' -gTA 3'	5' -gCA 3'	5' -TgA 3'	5' -TgA 3'	5' -TgA 3'	5' -TgA 3'	5' -TAT 3'	5' -gTA 3'
3'-primer(s) ³	78(319)	57(257)	57(257)	67(286)	57(256)	78(319)	86(344)	57(257)
	5' -CAC 3'	5' -CgA 3'	5' -CAT 3'	5' -gAA 3'	5' -gCT 3'	5' -gTA 3'	5' -CCA 3'	5' -CgA 3'
			57(258)		67(286)	87(348)		
			5' -gCg 3'		5' -gAg 3'	5' -CAT 3'		
Well No.	1	2	3	4	5	6	7	8
DRB1 allele ⁴								
*09:01:02-09:01:07	1	2						
*09:02:01-09:02:02	1		3					
*09:03	1	2		4				
*09:04	1	2			5			
*09:05		2				6		
*09:06	1	2					7	
*09:07	1	2						8
*09:08	1				5			
*09:09	1	2				6		
DRB1 allele ⁴								
Well No.	1	2	3	4	5	6	7	8